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# INFORMATION DISCLOSURE STATEMENT BY APPLICANT

(Use as many sheets as necessary)

## Complete if Known

Application Number	10/037,718
Filing Date	04 January 2002
First Named Inventor	McGinnis, R.E.
Art Unit	1631
Examiner Name	McGinnis, R.E.
Attorney Docket Number	2DLSM&R12/01

Sheet 6

of

10

## NON PATENT LITERATURE DOCUMENTS

Examiner Initials*	Cite No. <sup>1</sup>	Include name of the author (in CAPITAL LETTERS), title of the article (when appropriate), title of the item (book, magazine, journal, serial, symposium, catalog, etc.), date, page(s), volume-issue number(s), publisher, city and/or country where published.	T <sup>2</sup>
CA		DELAHUNTY&NICKERSON, ET AL, Am J Hum Genet. 1996 Jun;58(6):1239-46. Testing the feasibility of DNA typing for human identification by PCR and an oligonucleotide ligation assay	
CB		KWOK ET AL, Genomics. 1996 Jan 1;31(1):123-6. Increasing the information content of STS-based genome maps: identifying polymorphisms in mapped STSs	
CC		BUETOW ET AL, Nat Genet. 1999 Mar;21(3):323-5. Reliable identification of large numbers of candidate SNPs from public EST data	
CD		PICOULT-NEWBERG & NICKERSON, ET AL, Genome Res. 1999 9: 167-174, Mining SNPs From EST Databases	
CE		ROSES ET AL, Genomics. 1998 Nov 15;54(1):31-8. A 4-Mb high-density single nucleotide polymorphism-based map around human APOE.	
CF		TOBE & NICKERSON, ET AL, Nucleic Acids Res. 1996 Oct 1;24(19):3728-32. Single-well genotyping of diallelic sequence variations by a two-color ELISA-based oligonucleotide ligat	
CG		HACIA, FODOR, COLLINS, ET AL, Nucleic Acids Res. 1998 Nov 1;26(21):4975-82. Enhanced high density oligonucleotide array-based sequence analysis using modified nucleoside triph	
CH		KNAPP, ET AL, Nucleic Acids Res. 1994 Oct 11;22(20):4167-75. Genetic Bit Analysis: a solid phase method for typing single nucleotide polymorphisms	
CI		ZHAO, ET AL, Am J Hum Genet. 1998 Jul;63(1):225-40. Mapping of complex traits by single-nucleotide polymorphisms	
CJ		DERISI & BROWN, Science Vol. 278, No. 5338, (1997), pp. 680-686 Exploring the Metabolic and Genetic Control of Gene Expression on a Genomic Scale	

Examiner Signature	/Pablo Whaley/	Date Considered	10/13/2009
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\*EXAMINER: Initial if reference considered, whether or not citation is in conformance with MPEP 609. Draw line through citation if not in conformance and not considered. Include copy of this form with next communication to applicant.

<sup>1</sup> Applicant's unique citation designation number (optional). <sup>2</sup> Applicant is to place a check mark here if English language Translation is attached. This collection of information is required by 37 CFR 1.98. The information is required to obtain or retain a benefit by the public which is to file (and by the USPTO to process) an application. Confidentiality is governed by 35 U.S.C. 122 and 37 CFR 1.14. This collection is estimated to take 2 hours to complete, including gathering, preparing, and submitting the completed application form to the USPTO. Time will vary depending upon the individual case. Any comments on the amount of time you require to complete this form and/or suggestions for reducing this burden, should be sent to the Chief Information Officer, U.S. Patent and Trademark Office, P.O. Box 1450, Alexandria, VA 22313-1450. DO NOT SEND FEES OR COMPLETED FORMS TO THIS ADDRESS. SEND TO: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450.

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ALL REFERENCES CONSIDERED EXCEPT WHERE LINED THROUGH. /P.W./